

SEQUENCE LISTING

<110> Gish, Kurt C.

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TECH CENTER 1600/2900

Mack, David

<120> NOVEL METHODS OF DIAGNOSING BREAST CANCER, COMPOSITIONS, AND METHODS OF SCREENING FOR BREAST CANCER MODULATORS

<130> A-69026/DJB/JJD

<140> US 09/702,216

<141> 2000-10-30

<150> US 09/525,361

<151> 2000-03-15

<150> US 09/453,137

<151> 1999-12-02

<150> US 09/450,810

<151> 1999-11-29

<150> PCT/US 00/06952

<151> 2000-03-15

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<170> PatentIn Ver. 2.1

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<221> misc feature

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<223> "n" at positions 10118 and 10180 can be any base.

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Ser Asp Ala Ala Glu Leu Asn His Lys Glu Glu His Ser Leu His Val
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Gln Asp Pro Ser Ser Ser Lys Lys Asp Leu Lys Ser Ala Val Leu
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Ser Glu Lys Ala Gly Phe Asn Tyr Glu Ser Pro Ser Lys Gly Gly Asn 85 90 95

Phe Pro Ser Phe Pro His Asp Glu Val Thr Asp Arg Asn Met Leu Ala

Phe Ser Phe Pro Ala Ala Gly Gly Val Cys Glu Pro Leu Lys Ser Pro Gln Arg Ala Glu Ala Asp Asp Pro Gln Asp Met Ala Cys Thr Pro Ser Gly Asp Ser Leu Glu Thr Lys Glu Asp Gln Lys Met Ser Pro Lys Ala Thr Glu Glu Thr Gly Gln Ala Gln Ser Gly Gln Ala Asn Cys Gln Gly Leu Ser Pro Val Ser Val Ala Ser Lys Asn Pro Gln Val Pro Ser Asp Gly Gly Val Arg Leu Asn Lys Ser Lys Thr Asp Leu Leu Val Asn Asp Asn Pro Asp Pro Ala Pro Leu Ser Pro Glu Leu Gln Asp Phe Lys Cys Asn Ile Cys Gly Tyr Gly Tyr Tyr Gly Asn Asp Pro Thr Asp Leu Ile Lys His Phe Arg Lys Tyr His Leu Gly Leu His Asn Arg Thr Arg Gln Asp Ala Glu Leu Asp Ser Lys Ile Leu Ala Leu His Asn Met Val Gln Phe Ser His Ser Lys Asp Phe Gln Lys Val Asn Arg Ser Val Phe Ser

275 280 285

Gly Val Leu Gln Asp Ile Asn Ser Ser Arg Pro Val Leu Leu Asn Gly 290 295 300

Thr Tyr Asp Val Gln Val Thr Ser Gly Gly Thr Phe Ile Gly Ile Gly 305 310 315

Arg Lys Thr Pro Asp Cys Gln Gly Asn Thr Lys Tyr Phe Arg Cys Lys 325 330 335

Phe Cys Asn Phe Thr Tyr Met Gly Asn Ser Ser Thr Glu Leu Glu Gln 340 345 350

His Phe Leu Gln Thr His Pro Asn Lys Ile Lys Ala Ser Leu Pro Ser

Ser Glu Val Ala Lys Pro Ser Glu Lys Asn Ser Asn Lys Ser Ile Pro Ala Leu Gln Ser Ser Asp Ser Gly Asp Leu Gly Lys Trp Gln Asp Lys Ile Thr Val Lys Ala Gly Asp Asp Thr Pro Val Gly Tyr Ser Val Pro Ile Lys Pro Leu Asp Ser Ser Arg Gln Asn Gly Thr Glu Ala Thr Ser Tyr Tyr Trp Cys Lys Phe Cys Ser Phe Ser Cys Glu Ser Ser Ser Leu Lys Leu Leu Glu His Tyr Gly Lys Gln His Gly Ala Val Gln Ser Gly Gly Leu Asn Pro Glu Leu Asn Asp Lys Leu Ser Arg Gly Ser Val Ile Asn Gln Asn Asp Leu Ala Lys Ser Ser Glu Gly Glu Thr Met Thr Lys Thr Asp Lys Ser Ser Ser Gly Ala Lys Lys Asp Phe Ser Ser Lys Gly Ala Glu Asp Asn Met Val Thr Ser Tyr Asn Cys Gln Phe Cys Asp Phe Arg Tyr Ser Lys Ser His Gly Pro Asp Val Ile Val Val Gly Pro Leu Leu Arg His Tyr Gln Gln Leu His Asn Ile His Lys Cys Thr Ile Lys His Cys Pro Phe Cys Pro Arg Gly Leu Cys Ser Pro Glu Lys His Leu Gly Glu Ile Thr Tyr Pro Phe Ala Cys Arg Lys Ser Asn Cys Ser His Cys Ala Leu Leu Leu His Leu Ser Pro Gly Ala Ala Gly Ser Ser Arg Val Lys His Gln Cys His Gln Cys Ser Phe Thr Thr Pro

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865 870 875 880

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Ala Asn Cys Leu Thr Thr Lys Thr Ser Leu Trp Arg Lys Asn Ala Asn 900 905 910

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Gln Arg Glu Ile Pro Leu Pro Ser Leu Ser Lys Tyr Glu Ala Gln Gly 995 1000 1005

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1060 1065 1070

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Lys Tyr Gln Tyr Pro Leu Phe Gly Leu Pro Phe Val His Asn Asp Phe 1090 1095 1100

Gln Ser Glu Ala Asp Trp Leu Arg Phe Trp Ser Lys Tyr Lys Leu Ser 1105 1110 1115 1120

Val Pro Gly Asn Pro His Tyr Leu Ser His Val Pro Gly Leu Pro Asn

Pro Cys Gln Asn Tyr Val Pro Tyr Pro Thr Phe Asn Leu Pro Pro His 1140 1145 1150

Phe Ser Ala Val Gly Ser Asp Asn Asp Ile Pro Leu Asp Leu Ala Ile 1155 1160 1165

Lys His Ser Arg Pro Gly Pro Thr Ala Asn Gly Ala Ser Lys Glu Lys 1170 1180

Thr Lys Ala Pro Pro Asn Val Lys Asn Glu Gly Pro Leu Asn Val Val 1185 1190 1195 1200

Lys Thr Glu Lys Val Asp Arg Ser Thr Gln Asp Glu Leu Ser Thr Lys 1205 1210 1215

Cys Val His Cys Gly Ile Val Phe Leu Asp Glu Val Met Tyr Ala Leu 1220 1225 1230

His Met Ser Cys His Gly Asp Ser Gly Pro Phe Gln Cys Ser Ile Cys 1235 1240 1245

Gln His Leu Cys Thr Asp Lys Tyr Asp Phe Thr Thr His Ile Gln Arg 1250 1255 1260

Gly Leu His Arg Asn Asn Ala Gln Val Glu Lys Asn Gly Lys Pro Lys 1265 1270 1275 1280

Glu

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<213> Unknown Organism

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<223> "Xaa" at position 3 can be any base

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<223> Description of Unknown Organism: cytokine receptor extracellular motif found in many species

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Trp Ser Xaa Trp Ser